

# CRF Errors Corrected by the STIC Systems Branch

CRF Processing Date: 01/26/2002

Edited by: ALP

Verified by: ALP (STIC staff)

Serial Number: 10/053,530

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☒ Other: Seq 3 - inserted hard return

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



OIPE

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/053,530

DATE: 02/26/2002

TIME: 12:23:09

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Output Set: N:\CRF3\02262002\J053530.raw

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2   Hayden-Ledbetter, Martha
3 <120> TITLE OF INVENTION: Binding Domain-Immunoglobulin Fusion Proteins
4 <130> FILE REFERENCE: 390069.401
5 <140> CURRENT APPLICATION NUMBER: US/10/053,530
6 <141> CURRENT FILING DATE: 2002-01-17
7 <150> PRIOR APPLICATION NUMBER: US 09/765,208
8 <151> PRIOR FILING DATE: 2001-01-17
9 <160> NUMBER OF SEQ ID NOS: 38
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 PATENT APPLICATION: US/10/053,530

DATE: 02/26/2002  
 TIME: 12:23:09

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54 <221> NAME/KEY: C_region
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56 <223> OTHER INFORMATION: HUMAN IgG1 Fc TAIL, WILD TYPE HINGE, CH2 AND CH3
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95 <222> LOCATION: (808)..(1513)

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RAW SEQUENCE LISTING  
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96 <223> OTHER INFORMATION: HINGE CYSTEINES (826-829; 844-847; 853-856) MUTATED TO
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98   PROLINE TO SERINE MUTATION (880-883) IN CH2 DISRUPTS EFFECTOR FUN
99   CTION
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103   ccaggggaga aggtcacaat gacttgcagg gccagctcaa gtgtaagtta catgcactgg      180
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115   acatccccac cgtccccagc acctgaatc ctggggggat cgtcagctct cctcttcccc      900
116   ccaaaaccca aggacaccct catgatctcc cggaccctg aggtcacatg cgtgggtggtg      960
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137 <221> NAME/KEY: C_region
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PATENT APPLICATION: US/10/053,530

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188   gtcacaatga cttgcagggc cagctcaagt gtaagttaca tgcactggta ccagcagaag      180
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VERIFICATION SUMMARY

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Application Serial Number: US/10/053,530  
Alpha or Numeric: Numeric  
Application Class:  
Application File Date: 01-17-2002  
Art Unit: OIPE  
Software Application: PatentIN3.0  
Total Number of Sequences: 38  
Total Nucleotides: 14732  
Total Amino Acids: 4719  
Number of Errors: 0  
Number of Warnings: 0  
Number of Corrections: 0

MESSAGE SUMMARY